AMENDMENTS TO THE CLAIMS

1. (Currently amended) A compound of formula (1)

$$R_1-A-X-CH_2-R_3-R_4-L_1$$
 (1),

wherein:

A is a group recognized by O^6 alkylguanine DNA alkyltransferases (AGT) as a substrate the group R₁-A is a purine radical of formula (2)

$$R_8 \xrightarrow{N} N R_6 \qquad (2)$$

X is oxygen-or sulfur:

R₁ is a group -R₂-L₂ or a group R₅;

R₂ and R₄ are, independently of each other, a linker straight or branched chain alkylene group or polyvalent branched chain alkyl group with 1 to 300 carbon atoms, wherein optionally

- (a) one or more carbon atoms are replaced by oxygen;
- (b) one or more carbon atoms are replaced by nitrogen carrying a hydrogen atom, and the adjacent carbon atom is substituted by oxo;
- (c) one or more carbon atoms are replaced by oxygen, and the adjacent carbon atom is substituted by oxo;
- (d) the bond between two adjacent carbon atoms is a double or a triple bond;
- (e) one or more carbon atoms are replaced by a phenylene, a saturated or unsaturated cycloalkylene, a saturated or unsaturated bicycloalkylene, a bridging heteroaromatic or a bridging saturated or unsaturated heterocyclyl group;
- (f) two adjacent carbon atoms are replaced by a disulfide linkage;

or a combination of two or more alkylene and/or modified alkylene groups as defined under (a) to (f) above, optionally containing substituents;

 R_3 is an aromatic or a heteroaromatic group, or an optionally substituted <u>1-alkenylene</u>,

<u>1-alkynylene</u>, <u>1-cycloalkenylene</u>, or <u>an</u> unsaturated alkyl, cycloalkyl or heterocyclyl group with the double bond connected to CH₂;

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R₃ is arylmethyl or heteroarylmethyl or an optionally substituted cycloalkyl, cycloalkenyl or heterocyclyl group;

R6 is hydrogen, hydroxy or unsubstituted or substituted amino;

one of R7 and R8 is R1 and the other one is hydrogen; and

L₁ is a label, and L₂ are

one or a plurality of same or different labels selected from the group consisting of a spectroscopic probe, a magnetic probe, a contrast reagent, a radioactive moiety, avidin, streptavidin, biotin, a moiety which is capable of crosslinking to other molecules selected from the group consisting of a maleimide, active ester, azide and benzophenone, a tethered metalchelate which is capable of generating hydroxyl radicals upon exposure to H₂O₂ and ascorbate, malachite green, a moiety covalently attached to a solid support, a lipid, methotrexate, a linear poly(arginine) of D- and/or L-arginine with 6-15 arginine residues, a linear polymer of 6-15 subunits each carrying a guanidinium group, oligomers or short-length polymers of 6-50 subunits, a portion of which have attached guanidinium groups, and parts of a sequence of a HIV-tat protein;

or L_{1, is.}-a bond connecting R₄ to A forming a cyclic substrate, e-a further group -R₃-CH₂-X-A-R₁, or a nucleic acid or a derivative thereof capable of undergoing base-pairing with its complementary strand; end-or

L₂ is a <u>nucleic acid or a derivative thereof capable of undergoing base-pairing with its</u> complementary strand if R₂ is hydrogen tabel or a plurality of same or different tabels.

(Currently amended) The compound according to claim 1, of formula (1) wherein
 A is a heteroaromatic group containing 1 to 5 nitrogen atoms;

X is oxygen;

R₊ is a group - R₂-L₂ or a group R₅;

R₂ and R₄ are, independently of each other, a straight or branched chain alkylene group with 1 to 300 earbon atoms, wherein optionally

(a) one or more carbon atoms are replaced by oxygen, in particular wherein every third carbon atom is replaced by oxygen, e.g. a poylethyleneoxy group with 1 to 100 ethyleneoxy units;

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(b) one or more carbon atoms are replaced by nitrogen carrying a hydrogen atom, and the adjacent carbon atoms are substituted by oxo, representing an amide function

NH_CO :

- (e) one or more carbon atoms are replaced by oxygen, and the adjacent carbon atoms are substituted by oxo, representing an ester function O-CO;
- (d) the bond between two adjacent earbon atoms is a double or a triple bond, representing a function CH=CH or CEC;
- (e) one or more carbon atoms are replaced by a phenylene, a saturated or unsaturated eyeloalkylene, a saturated or unsaturated bicycloalkylene, a bridging heteroaromatic or a bridging saturated or unsaturated heteroeyelyl group;
- (f) two adjacent carbon atoms are replaced by a disulfide linkage S-S-;
 or a combination of two or more, especially two or three, alkylene and/or modified alkylene
 groups as defined under (a) to (f) hereinbefore, optionally containing substituents;

R₃ is phenylphenylene, an unsubstituted or substituted mono- or bicyclic bridging heteroaryl group of 5 or 6 rings atoms comprising zero, one, two, three or four ring nitrogen atoms and zero or one oxygen atom and zero or one sulfur atom, with the proviso that at least one ring carbon atom is replaced by a nitrogen, oxygen or sulfur atom, 1-alkenyl, 1-alkinyl, 1-eyelohexenyl 1-alkenylene, 1-alkinylene, 1-cyclohexenylene, with 3 to 7 carbon atoms, wherein the double or triple bond is connected to CH_{2x} or an optionally substituted unsaturated bridging heterocyclyl group with 3 to 12 atoms and 1 to 5 heteroatoms selected from the group consisting of nitrogen, oxygen and sulfur, and a double bond in the position connecting the heterocyclyl group to methylene-CH₃; and

R₃ is optionally substituted phenylmethyl or naphthylmethyl; optionally substituted heteroarylmethyl wherein heteroaryl is a mono—or bicyclic heteroaryl group comprising zero, one, two, three or four ring nitrogen atoms and zero or one oxygen atom and zero or one sulfur atom, with the proviso that at least one ring carbon atom is replaced by a nitrogen, oxygen or sulfur atom, and which has 5 to 12 ring atoms; optionally substituted cycloalkyl with 3 to 7 carbon atoms; optionally substituted cycloalkenyl with 5 to 7 carbon atoms; or optionally

substituted saturated or unsaturated heterocyclyl with 3 to 12 atoms, and 1 to 5 heteroatoms selected from the group consisting of nitrogen, oxygen and sulfur-

 L_i is one or a plurality of same or different labels selected from a spectroscopic probe, a magnetic probe, a contrast reagent, a molecule which is one part of a specific binding pair which is capable of specifically binding to a partner, a molecule that is suspected to interact with other biomolecules, a library of molecules that are suspected to interact with other biomolecules, a molecule which is capable of crosslinking to other molecules, a molecule which is capable of generating hydroxyl radicals upon exposure to H_2O_2 and ascorbate, a molecule which is capable of generating reactive radicals upon irradiation with light, a molecule covalently attached to a solid support, a nucleic acid or a derivative thereof capable of undergoing base pairing with its complementary strand, a lipid or other hydrophobic molecule with membrane inserting properties, a biomolecule with desirable enzymatic, chemical or physical properties, a bond connecting R_4 to Λ forming a cyclic substrate, and a further group R_2 CH_2 X Λ R_4 ; and

L₂ is one or a plurality of same or different labels selected from a spectroscopic probe, a magnetic probe, a contrast reagent, a molecule which is one part of a specific binding pair which is capable of specifically binding to a partner, a molecule that is suspected to interact with other biomolecules, a library of molecules that are suspected to interact with other biomolecules, a molecule which is capable of crosslinking to other molecules, a molecule which is capable of generating hydroxyl radicals upon exposure to H₂O₂ and ascorbate, a molecule which is capable of generating reactive radicals upon irradiation with light, a molecule covalently attached to a solid support, a lipid or other hydrophobic molecule with membrane inserting properties, and a biomolecule with desirable enzymatic, chemical or physical properties.

3. (Cancelled)

 (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein X is oxygen and R₂ is phenyl R₃ is phenylene.

- (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein X is oxygen and R₂ is thienyl R₃ is thicnylene.
- 6. (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein the group R₁. A is a purine radical of formula (2), R₆ is unsubstituted amino, R₇ is R₁, and R₈ is hydrogen, and X is oxygen.
- 7. (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein the group R₄ A is a purine radical of formula (2), R₆ is unsubstituted amino, R₇ is a group -R₂-L₂, and R₈ is hydrogen, and X is oxygen.
- (Currently amended) The compound according to claim 7₁ wherein L₂ is a spectroscopic probe.
- (Currently amended) The compound according to claim 7₂ wherein L₁ and L₂ are spectroscopic probes.
- 10. (Currently amended) The compound according to claim 9_a wherein L_1 and L_2 represent a fluorescence donor / fluorescence quencher pair.
- 11. (Currently amended) The compound according to claim 10, wherein L1 and L2 represent a FRET pair.
- 12. (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein the group R₁. A is a purine radical of formula (2), R₆ is unsubstituted amino, R₇ is a group R₅, and R₈ is hydrogen, and X is oxygen.
- 13. (Currently amended) The compound according to claim 12, wherein R5 is cyclopentyl.

- 14. (Currently amended) The compound according to elaim 3 of formula (1) claim 1, wherein the group R₁. A is a purine radical of formula (2), R₆ is unsubstituted amino, R₇ is hydrogen, and R₈ is R₁₇ and X is oxygen.
- 15. (Currently amended) The compound according to claim 3 of formula (1) claim 1, wherein the group R₁—A is a purine radical of formula (2), R₆ is unsubstituted amino, R₇ is hydrogen, and R₈ is a group -R₂-L₂, and X is oxygen.
- 16. (Currently amended) The compound according to claim 15, wherein L₂ is a spectroscopic probe.
- 17. (Currently amended) The compound according to claim 15_x wherein L₁ and L₂ are spectroscopic probes.
- 18. (Currently amended) The compound according to claim 17, wherein L₁ and L₂ represent a fluorescence donor / fluorescence quencher pair.
- 19. (Currently amended) The compound according to claim 18, wherein L₁ and L₂ represent a FRET pair.
- (Currently amended) The compound according to claim 15, wherein L2 is <u>avidin</u>, <u>streptavidin or biotina molecule representing one part of a specific binding pair</u>.
- (Currently amended) The compound according to claim 15, wherein L₂ is a moiety molecule-covalently attached to a solid support.
- 22. (Currently amended) The compound according to claim 15, wherein L2 is a linear poly(arginine) of D- and/or L-arginine with 6-15 arginine residues, a linear polymer of 6-15 subunits each carrying a guanidinium group, oligomers or short-length polymers of 6-50 subunits, a portion of which have attached guanidinium groups, or parts of a sequence of a HIV-tat protein-cell membrane transport enhancer group.

23-43. (Cancelled)

- 44. (Currently amended) A method for detecting and/or manipulating a protein of interest, wherein the protein of interest is incorporated into-fused to an AGT-fusion protein, the AGT fusion protein is contacted with a compound of formula (1) according to claim 1, and the AGT fusion protein is detected and optionally further manipulated using the label L₁ in a system designed for recognizing and/or handling the label.
- 45. (Original) The method according to claim 44, wherein in the compound of formula (1) label L₂ is a solid support, and the AGT fusion protein contacted with the compound of formula (1) is separated from the compound of formula (1) by filtration or centrifugation or separation of magnetic beads.
- 46. (Currently amended) The method according to claim 44, wherein in the compound of formula (1) label L₁ is one member and label L₂ the other member of two interacting spectroscopic probes L₁ / L₂, wherein energy can be transferred nonradiatively through dynamic or static quenching, and the AGT fusion protein is detected by fluorescence.
- 47. (Currently amended) The method according to claim 44 for detecting and/or manipulating a protein of interest, wherein the protein of interest is fused with a mutant AGT, the mutant AGT fusion protein is contacted with a mixture of
- (a) a compound of formula (1) wherein R₁ is a group R₅, and which is not recognized by does not react with the mutant AGT, and
- (b) another compound of formula (1)_a recognized by which reacts with the mutant AGT fusion protein,

and the mutant AGT fusion protein is detected and optionally further manipulated using the label in a system designed for recognizing and/or handling the label.